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25. (new) A variant according to claim 23 wherein the amino acid mutation is in the B domain and/or the C domain of the HBV DNA polymerase.

26. (new) A variant according to claim 23 comprising a mutation in one or more of amino acids within the sequence:

Q/K T Y/F G R/W KLHL Y/L S/A HPI I/V LGFRK I/L PMG V/G GLS PFLL

AQFTSAI C/L S (SEQ ID NO:25)

of HBV DNA polymerase.

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27. (new) A variant according to claim 26 comprising a mutation in one or more amino acids within the sequence:

S/A HPI I/V LGFRK I/L PMG V/G GLSPFLLAQFTSAIC/L S (SEQ ID NO:44)

of HBV DNA polymerase.

28. (new) A variant according to claim 23 comprising a nucleotide sequence which encodes a DNA polymerase having the amino acid sequence:

X<sub>1</sub>HPIX<sub>2</sub>LGX<sub>3</sub>RKX<sub>4</sub>PMGX<sub>5</sub>GLSX<sub>6</sub>FLX<sub>7</sub>AQFTSAX<sub>8</sub>X<sub>9</sub> ..... (SEQ ID NO:27)

X<sub>10</sub>FX<sub>11</sub>YX<sub>12</sub>DDX<sub>13</sub>VLGAX<sub>14</sub>X<sub>15</sub> (SEQ ID NO:28)

wherein X<sub>1</sub> is S or A;

X<sub>2</sub> is I or V;

X<sub>3</sub> is F or L;

X<sub>4</sub> is I or L;

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X<sub>5</sub> is L or V or G;

X<sub>6</sub> is P or L;

X<sub>7</sub> is L or M;

X<sub>8</sub> is I or L;

X<sub>9</sub> is C or L;

X<sub>10</sub> is A or V;

X<sub>11</sub> is S or A;

X<sub>12</sub> is M or I or V;

X<sub>13</sub> is V or L or M;

X<sub>14</sub> is K or R; and/or

X<sub>15</sub> S or T;

and wherein said variant exhibits reduced sensitivity to a nucleoside analogue, such as famciclovir (penciclovir) and/or lamivudine (3TC).

29. (new) A variant according to claim 23 having a mutation in one or more of amino acids 118 to 169 or 169 to 207 of HBV surface antigen.

30. (new) A variant according to claim 29 comprising a DNA polymerase having the amino acid sequence:

X<sub>16</sub>TX<sub>17</sub>X<sub>18</sub>X<sub>19</sub>KLHLX<sub>20</sub>X<sub>21</sub>HPIX<sub>22</sub>LGX<sub>3</sub>RKX<sub>4</sub>PMGX<sub>5</sub>GLSX<sub>6</sub>FLX<sub>7</sub>AQFTSAX<sub>8</sub>X<sub>9</sub>.....

(SEQ ID NO:42)

X<sub>10</sub>FX<sub>11</sub>YM<sub>12</sub>DDX<sub>13</sub>VLGAX<sub>14</sub>X<sub>15</sub> (SEQ ID NO:43)

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wherein: X<sub>16</sub> is Q or K;  
X<sub>17</sub> is Y or F;  
X<sub>18</sub> is G;  
X<sub>19</sub> is R or W or E;  
X<sub>20</sub> is Y or L;  
X<sub>21</sub> is S or A;  
X<sub>22</sub> is I or V;  
X<sub>3</sub> is F or L;  
X<sub>4</sub> is I or L;  
X<sub>5</sub> is L or V or G;  
X<sub>6</sub> is P or L;  
X<sub>7</sub> is L or M;  
X<sub>8</sub> is I or L;  
X<sub>9</sub> is C or L;  
X<sub>10</sub> is A or V;  
X<sub>11</sub> is S or A;  
X<sub>12</sub> is M or I or V;  
X<sub>13</sub> is V or L or M;  
X<sub>14</sub> is K or R; and/or  
X<sub>15</sub> S or T;

and wherein said variant exhibits reduced sensitivity to a nucleoside analogue, such as famciclovir (penciclovir) and/or lamivudine (3TC).